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DATE: 03/15/2001

#2

RAW SEQUENCE LISTING PATENT APPLICATION US/09/726,258

TIME: 22:58:34

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This Raw Listing contains the General Information Section and up to the first 5 pages.

ENTERED SEQUENCE LISTING 1 2 3 (1) General Information: 4 (i) APPLICANT: Genentech, Inc., Hsei, Vanessa 5 6 Koumenis, Iphigenia 7 Leong, Steven R. 8 Presta, Leonard G. 9 Shahrokh, Zahra 10 Zapata, Gerardo A. 11 (ii) TITLE OF INVENTION: ANTIBODY FRAGMENT-POLYMER CONJUGATES AND 12 13 HUMANIZED ANTI-IL-8 MONOCLONAL ANTIBODIES 14 (iii) NUMBER OF SEQUENCES: 72 15 16 17 (iv) CORRESPONDENCE ADDRESS: (A) ADDRESSEE: Genentech, Inc. 18 (B) STREET: 1 DNA Way 19 (C) CITY: South San Francisco 20 21 (D) STATE: California 22 (E) COUNTRY: USA (F) ZIP: 94080 23 25 (v) COMPUTER READABLE FORM: 26 (A) MEDIUM TYPE: 3.5 inch, 1.44 Mb floppy disk 27 (B) COMPUTER: IBM PC compatible 28 (C) OPERATING SYSTEM: PC-DOS/MS-DOS (D) SOFTWARE: WinPatin (Genentech) 29 30 (vi) CURRENT APPLICATION DATA: 31 32 (A) APPLICATION NUMBER: 09/726,258 33 (B) FILING DATE: 34 (C) CLASSIFICATION: 35 (vii) PRIOR APPLICATION DATA: 36 37 (A) APPLICATION NUMBER: 09/234,182 (B) FILING DATE: 38 39 40 (vii) PRIOR APPLICATION DATA: (A) APPLICATION NUMBER: 60/094003 41 42 (B) FILING DATE: 24-JUL-1998 43 44 45 (viii) ATTORNEY/AGENT INFORMATION:

(A) NAME: Love, Richard B.

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RAW SEQUENCE LISTING PATENT APPLICATION US/09/726,258

DATE: 03/15/2001 TIME: 22:58:34

47		(B) REGISTRATION NUMBER: 34,659 (C) REFERENCE/DOCKET NUMBER: P1085R4-1A
48		(C) REFERENCE/DOCKET NUMBER: PIUGSK4-IA
49		THE PROPERTY OF THE PARTY OF TH
50	(ix)	TELECOMMUNICATION INFORMATION:
51		(A) TELEPHONE: 650/225-5530
52		(B) TELEFAX: 650/952-9881
53	(2) INI	FORMATION FOR SEQ ID NO:1:
54		
55	(i)	SEQUENCE CHARACTERISTICS:
56		(A) LENGTH: 22 base pairs
57		(B) TYPE: Nucleic Acid
58		(C) STRANDEDNESS: Single
59		(D) TOPOLOGY: Linear
60		
61	(xi)	SEQUENCE DESCRIPTION: SEQ ID NO:1:
62		
63		
64	CAGTC	CAACT GTTCAGGACG CC 22
65		
66	(2) IN	FORMATION FOR SEQ ID NO:2:
67		
68	(i)	SEQUENCE CHARACTERISTICS:
69		(A) LENGTH: 22 base pairs
70		(B) TYPE: Nucleic Acid
71		(C) STRANDEDNESS: Single
72		(D) TOPOLOGY: Linear
73		
74	(xi)	SEQUENCE DESCRIPTION: SEQ ID NO:2:
75		
76		
77	GTGCT	GCTCA TGCTGTAGGT GC 22
78		
79	(2) IN	FORMATION FOR SEQ ID NO:3:
80		
81	(i)	SEQUENCE CHARACTERISTICS:
82		(A) LENGTH: 23 base pairs
83		(B) TYPE: Nucleic Acid
84		(C) STRANDEDNESS: Single
85		(D) TOPOLOGY: Linear
86		
87	(xi)	SEQUENCE DESCRIPTION: SEQ ID NO:3:
88		
89		
90	GAAGT'	IGATG TCTTGTGAGT GGC 23
91	(0)	CODMITTON FOR CHO ID NO.4.
92	(2) IN	FORMATION FOR SEQ ID NO:4:
93	123	GEOLIENGE CHARACTERICATOS.
94	(1)	SEQUENCE CHARACTERISTICS: (A) LENGTH: 24 base pairs
95		(A) mypr. Musleig Acid
96		(B) TYPE: Nucleic Acid (C) STRANDEDNESS: Single
97		
98		(D) TOPOLOGY: Linear
99		

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100	(xi) SEQUENCE DESCRIPTION: SEQ ID NO:4:
101	
102	
103	GCATCCTAGA GTCACCGAGG AGCC 24
104	
105	(2) INFORMATION FOR SEQ ID NO:5:
106	1-7
107	(i) SEQUENCE CHARACTERISTICS:
108	(A) LENGTH: 22 base pairs
109	(B) TYPE: Nucleic Acid
110	(C) STRANDEDNESS: Single
	(D) TOPOLOGY: Linear
111	(D) TOPOLOGI. Hincar
112	(with another programmer), SEO ID NO.5.
113	(xi) SEQUENCE DESCRIPTION: SEQ ID NO:5:
114	
115	
116	CACTGGCTCA GGGAAATAAC CC 22
117	
118	(2) INFORMATION FOR SEQ ID NO:6:
119	
120	(i) SEQUENCE CHARACTERISTICS:
121	(A) LENGTH: 22 base pairs
122	(B) TYPE: Nucleic Acid
123	(C) STRANDEDNESS: Single
124	(D) TOPOLOGY: Linear
125	1- /
	(xi) SEQUENCE DESCRIPTION: SEQ ID NO:6:
127	(351) 52 2 511 51
128	
	GGAGAGCTGG GAAGGTGTGC AC 22
130	GGAGAGCIGG GAAGGIGIGG 110 12
	(2) INFORMATION FOR SEQ ID NO:7:
131	(2) INFORMATION FOR BEQ 12 No
132	(i) SEQUENCE CHARACTERISTICS:
133	(A) LENGTH: 35 base pairs
134	(A) DENGIH: 35 Dase paris
135	(B) TYPE: Nucleic Acid
136	(C) STRANDEDNESS: Single
137	(D) TOPOLOGY: Linear
138	
139	(xi) SEQUENCE DESCRIPTION: SEQ ID NO:7:
140	
141	
142	ACAAACGCGT ACGCTGACAT CGTCATGACC CAGTC 35
143	
144	(2) INFORMATION FOR SEQ ID NO:8:
145	
146	(i) SEQUENCE CHARACTERISTICS:
147	(A) LENGTH: 35 base pairs
148	(B) TYPE: Nucleic Acid
149	(C) STRANDEDNESS: Single
150	(D) TOPOLOGY: Linear
	(D) 10101001. 1111001
151	(xi) SEQUENCE DESCRIPTION: SEQ ID NO:8:
152	(XI) PEONEMED DESCRIPTION: SEQ ID NO.0.



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154	
155	ACAAACGCGT ACGCTGATAT TGTCATGACT CAGTC 35
156	
157	(2) INFORMATION FOR SEQ ID NO:9:
158	
159	(i) SEQUENCE CHARACTERISTICS:
160	(A) LENGTH: 35 base pairs
161	(B) TYPE: Nucleic Acid
162	(C) STRANDEDNESS: Single
163	(D) TOPOLOGY: Linear
164	(D) TOPOBOGI. Bilical
165	(xi) SEQUENCE DESCRIPTION: SEQ ID NO:9:
166	(XI) SEQUENCE DESCRIPTION: SEQ ID NO.3.
167	
	A CARA COCOM A COCOMO A CAR COCOMO A CACOMO A CA
168	ACAAACGCGT ACGCTGACAT CGTCATGACA CAGTC 35
169	(a) TATIONNAMION FOR SEC ID NO 10
170	(2) INFORMATION FOR SEQ ID NO:10:
171	(1)
172	(i) SEQUENCE CHARACTERISTICS:
173	(A) LENGTH: 37 base pairs
174	(B) TYPE: Nucleic Acid
175	(C) STRANDEDNESS: Single
176	(D) TOPOLOGY: Linear
177	
178	(xi) SEQUENCE DESCRIPTION: SEQ ID NO:10:
179	
180	
181	GCTCTTCGAA TGGTGGGAAG ATGGATACAG TTGGTGC 37
182	
183	(2) INFORMATION FOR SEQ ID NO:11:
184	
185	(i) SEQUENCE CHARACTERISTICS:
186	(A) LENGTH: 39 base pairs
187	(B) TYPE: Nucleic Acid
188	(C) STRANDEDNESS: Single
189	(D) TOPOLOGY: Linear
190	
191	(xi) SEQUENCE DESCRIPTION: SEQ ID NO:11:
192	
193	
194	CGATGGGCCC GGATAGACCG ATGGGGCTGT TGTTTTGGC 39
195	
196	(2) INFORMATION FOR SEQ ID NO:12:
197	• • • • • • • • • • • • • • • • • • • •
198	(i) SEQUENCE CHARACTERISTICS:
199	(A) LENGTH: 39 base pairs
200	(B) TYPE: Nucleic Acid
201	(C) STRANDEDNESS: Single
202	(D) TOPOLOGY: Linear
203	(D) IOIOMOOI. MAILOUI
203	(xi) SEQUENCE DESCRIPTION: SEQ ID NO:12:
205	(AL) DEGOLITED DEDOCATE ITOM. DEG ID NO.18.



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206	
207	CGATGGGCCC GGATAGACTG ATGGGGCTGT CGTTTTGGC 39
208	
	(2) INFORMATION FOR SEQ ID NO:13:
210	(') CHONDIGE CHARACTERICATION
211	(i) SEQUENCE CHARACTERISTICS:
212	(A) LENGTH: 39 base pairs (B) TYPE: Nucleic Acid
213	(C) STRANDEDNESS: Single
214	(C) STRANDEDNESS: SINGLE (D) TOPOLOGY: Linear
215	(D) TOPOLOGI: Linear
216 217	(xi) SEQUENCE DESCRIPTION: SEQ ID NO:13:
217	(XI) SEQUENCE DESCRIPTION. BDQ IS NO.13.
219	
220	CGATGGGCCC GGATAGACGG ATGGGGCTGT TGTTTTGGC 39
221	CGMIGGGCCC GGMIMOMCGG MIGGGGCTGI TGTTITGGG GG
	(2) INFORMATION FOR SEQ ID NO:14:
223	(2) Intoldation for page 13 hours
224	(i) SEQUENCE CHARACTERISTICS:
225	(A) LENGTH: 39 base pairs
226	(B) TYPE: Nucleic Acid
227	(C) STRANDEDNESS: Single
228	(D) TOPOLOGY: Linear
229	, ,
230	(xi) SEQUENCE DESCRIPTION: SEQ ID NO:14:
231	· · · · ·
232	
233	CGATGGGCCC GGATAGACAG ATGGGGCTGT TGTTTTGGC 39
234	
235	(2) INFORMATION FOR SEQ ID NO:15:
236	
237	(i) SEQUENCE CHARACTERISTICS:
238	(A) LENGTH: 39 base pairs
239	(B) TYPE: Nucleic Acid
240	(C) STRANDEDNESS: Single
241	(D) TOPOLOGY: Linear
242	()) growing proprietary die ID Mo.15.
243	(xi) SEQUENCE DESCRIPTION: SEQ ID NO:15:
244	
245	CGATGGGCCC GGATAGACTG ATGGGGCTGT TGTTTTGGC 39
246	CGATGGGCCC GGATAGACTG ATGGGGCTGT TGTTTTGGC 39
247	(2) INFORMATION FOR SEQ ID NO:16:
	(2) INFORMATION FOR BEQ 1D NO.10.
249 250	(i) SEQUENCE CHARACTERISTICS:
251	(A) LENGTH: 369 base pairs
252	(B) TYPE: Nucleic Acid
253	(C) STRANDEDNESS: Double
254	(D) TOPOLOGY: Linear
255	• •
256	(xi) SEQUENCE DESCRIPTION: SEQ ID NO:16:
257	



SEQUENCE VERIFICATION REPORT PATENT APPLICATION US/09/726,258

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